SEQUENCE LISTING

<110> van der Kuyl, Antoinette C.
Cornelissen, Marion

<120> MEANS AND METHODS FOR TREATMENT EVALUATION

<130> 5244US (REN/P55190US00)

<140> 10/055,728
<141> 2002-01-23

<150> 60/325,722
<151> 2001-09-28

<150> EP 0120373.2
<151> 2001-09-28

<150> EP 01200228.3
<151> 2001-01-23

<160> 156

<170> PatentIn version 3.1

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<223> TAG004 (EST AI217565, GenBank number BE466728)

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tttcctgcct caggaactcc aataaaacat tttccatcca ac 102

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<223> TAG007 (keratin 14, GenBank number XM_008578)

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ggaagatccc ctctcctgcc caagcacttc acagctggac cctgcttcac cctcaccccc 180

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aa 242

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gacatcgtca gcatcctgga caagctcatt cgtgcccctg actccctcaa gaccctggct 120

gactttgacc cccgcgtgtc tatccggctc cccagcacga gcggctcgga gggggtgccc 180

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caaatacctc aatttttgtt tttgagtact gtactatcct gtaaatatat cttaagcagg 180

tttgttttca gcactgatgg aaaataccag tgttgggttt ttttttagtt gccaacagtt 240

gtatgtttgc tgattattta tgacctgaaa taatatattt cttcttctaa gaagacattt 300

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<210> 69
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<223> TAG012 (IL1 receptor, type 1, GenBank number XM_002686)

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gtt 63

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<223> TAG013 (ephrin B1, GenBank numbers XM_002535, BC002524)

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cctacctacc ccttttctct tggccagggg cctcgtatcc taccttttcct tgtcccctgg 120

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catttagtat tttgctcaaa gtctaaggga cc 212

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ttgcaggggt tggggaggga gacatccagc ttgggctttc ccctttggaa taaaccattg 180

gtctgtcaca aaaaaaaaaa aaaaaaaaaa aaaa 214

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ataagctcca aaaaaaaaaa aaaaaaaaaa aaa 93

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caaatcttta ttgaatggtt ttgctcagca ccacctttta atatattggc agtacttatt 120

atataaaagg taaaccagca ttctcaaaaa aaaaaaaaaa aaa 163

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acctgtgggc tgcttcacat ccaccccggc tccccctgcc agcaacgctc actctacccc 120

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atgagtaaat aaactcttca aggcc 205

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gagcgcgggc cgggcgttcc tttccagcgc gggcagccct tcgaggtgct catcatcgcg 120

tcagacgacg gcttcaaggc cgtggttggg gacgcccagt accaccactt ccgcc 175

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<220>
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cctgaggcat tatttgaccg agtctacact caccagagtg acgtctggtc ctttggggtc 120

ctgctctggg agatcttcac gctgggggc tccccgtacc ccggcatccc tgtggaggag 180

ctcttcaagc tgctgaagga gggc 204

<210> 77
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gagaccccag gaacaataaa atgtcttctc ccacc 95

<210> 78
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<212> DNA
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aaaaaaaaaa aaaaaaaaa 139

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gataggtcat tatgattttt taccatttcg acttacataa tgaaaaccaa ttcattttaa 120

atatcagatt attattttgt aagttgtgga aaaagctaat tgtagttttc attatgaagt 180

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tcctctggca gcatagatcc ttaggtgctc aataaaggtg tgctgtattg aaaaaaaaaa 180

aaaaaaaaaa aaaa 194

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catcctatga cgcaaaaaaa aaaaaaaaaa aaaaaaa 97

<210> 82
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gatgtacttt aaggtgtctt tatgaagttt gctattctgg caataaactt ttagacttta 120

aaaaaaaaaa aaaaaaaaaa aaa 143

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<210> 85
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aggagaccaa aggtcgctac tgca 24

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